SCORE Search Results Details for Application 10552515 and Search Result 20081001_124607_us-10-552-515-1.rapbm.

This page gives you Search Results detail for the Application 10552515 and Search Result 20081001_124607_us-10-552-515-1.rapbm.

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                         GenCore version 6.2.1
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OM protein - protein search, using sw model

Run on:         October  1, 2008, 12:46:15 ; Search time 829 Seconds
                                           (without alignments)
                                           1085.529 Million cell updates/sec

Title:          US-10-552-515-1
Perfect score:  4950
Sequence:       1 MRMAATAWAGLQGPPLPTLC.........SELSSHWTPFTVPKASQLQQ 933

Scoring table:  BLOSUM62
                Gapop 10.0 , Gapext 0.5

Searched:       4190237 seqs, 964526986 residues

Total number of hits satisfying chosen parameters:      4190237

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :      Published_Applications_AA_Main:*
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       Pred. No. is the number of results predicted by chance to have a
       score greater than or equal to the score of the result being printed,
       and is derived by analysis of the total score distribution.
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SUMMARIES

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Result           Query
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     1    4950   100.0    933   5   US-10-552-515-1           Sequence 1, Appli
     2   4531.5   91.5    885   7   US-11-599-845A-700        Sequence 700, App
     3   4364.5   88.2    843   7   US-11-599-845A-698        Sequence 698, App
     4    3736    75.5    898   5   US-10-450-763-45847       Sequence 45847, A
     5   1531.5   30.9    920   4   US-10-104-047-2574        Sequence 2574, Ap
     6   1531.5   30.9    920   6   US-11-072-512-2574        Sequence 2574, Ap
     7   1527.5   30.9    981   6   US-11-582-861-10841       Sequence 10841, A
     8    1511    30.5   1046   6   US-11-582-861-9875        Sequence 9875, Ap
     9    1508    30.5    981   6   US-11-443-428A-801153     Sequence 801153,
    10    1488    30.1    960   6   US-11-177-894-7           Sequence 7, Appli
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    17   1402.5   28.3    910   5   US-10-484-148-14          Sequence 14, Appl
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ALIGNMENTS

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US-10-552-515-1
; Sequence 1, Application US/10552515
; Publication No. US20060194204A1
; GENERAL INFORMATION:
;  APPLICANT: The Government of the United States of America as
;  APPLICANT:  represented by the Secretary of the Department of Health and
;  APPLICANT:  Human Services
;  APPLICANT:  Bera, Tapan K.
;  APPLICANT:  Pastan, Ira H.
;  APPLICANT:  Lee, Byungkook
;  TITLE OF INVENTION: GENE EXPRESSED IN PROSTATE CANCER AND METHODS OF USE
;  FILE REFERENCE: 4239-68223-02
;  CURRENT APPLICATION NUMBER: US/10/552,515
;  CURRENT FILING DATE:  2005-10-06
;  PRIOR APPLICATION NUMBER: PCT/US2004/10588
;  PRIOR FILING DATE: 2004-04-05
;  PRIOR APPLICATION NUMBER: 60/461,399
;  PRIOR FILING DATE: 2003-04-08
;  NUMBER OF SEQ ID NOS: 12
;  SOFTWARE: PatentIn version 3.2
; SEQ ID NO 1
;   LENGTH: 933
;   TYPE: PRT
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Splice Variant-Novel Gene Expressed in Prostate
US-10-552-515-1

  Query Match              100.0%;  Score 4950;  DB 5;  Length 933;
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  Matches  933;  Conservative   0;  Mismatches   0;  Indels   0;  Gaps   0;

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RESULT 2
US-11-599-845A-700
; Sequence 700, Application US/11599845A
; Publication No. US20080025981A1
; GENERAL INFORMATION:
;  APPLICANT: Young, Paul E.
;  APPLICANT:  Ebner, Reinhard
;  APPLICANT:  Weaver, Zoe
;  APPLICANT:  Strovel, Jeffrey W.
```

```
;   APPLICANT:  Horrigan, Stephen K.
;   APPLICANT:  Shea, Martin
;   APPLICANT:  Weigle, Bernd
;   APPLICANT:  Rieger, Michael
;   APPLICANT:  Rick, Jennifer A.
;   APPLICANT:  Cain, Colyn B.
;   TITLE OF INVENTION: Cancer-linked Genes as Target for Chemotherapy
;   FILE REFERENCE: 689290-273
;   CURRENT APPLICATION NUMBER: US/11/599,845A
;   CURRENT FILING DATE:  2006-11-15
;   PRIOR APPLICATION NUMBER: 10/585,466
;   PRIOR FILING DATE: 2005-01-04
;   PRIOR APPLICATION NUMBER: PCT/US2005/000040
;   PRIOR FILING DATE: 2005-01-04
;   PRIOR APPLICATION NUMBER: 10/583,832
;   PRIOR FILING DATE: 2004-12-16
;   PRIOR APPLICATION NUMBER: PCT/US2004/42406
;   PRIOR FILING DATE: 2004-12-16
;   PRIOR APPLICATION NUMBER: 10/575,337
;   PRIOR FILING DATE: 2004-10-07
;   PRIOR APPLICATION NUMBER: PCT/US2004/33072
;   PRIOR FILING DATE: 2004-10-07
;   PRIOR APPLICATION NUMBER: 10/540,310
;   PRIOR FILING DATE: 2003-12-19
;   PRIOR APPLICATION NUMBER: PCT/US2003/40710
;   PRIOR FILING DATE: 2003-12-19
;   PRIOR APPLICATION NUMBER: 10/518,039
;   PRIOR FILING DATE: 2003-06-10
;   PRIOR APPLICATION NUMBER: PCT/US2003/19741
;   PRIOR FILING DATE: 2003-06-10
;   Remaining Prior Application data removed - See File Wrapper or PALM.
;   NUMBER OF SEQ ID NOS: 769
;   SOFTWARE: PatentIn version 3.0
; SEQ ID NO 700
;   LENGTH: 885
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-599-845A-700

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RESULT 3
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; Sequence 698, Application US/11599845A
; Publication No. US20080025981A1
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; GENERAL INFORMATION:
;  APPLICANT:  Young, Paul E.
;  APPLICANT:  Ebner, Reinhard
;  APPLICANT:  Weaver, Zoe
;  APPLICANT:  Strovel, Jeffrey W.
;  APPLICANT:  Horrigan, Stephen K.
;  APPLICANT:  Shea, Martin
;  APPLICANT:  Weigle, Bernd
;  APPLICANT:  Rieger, Michael
;  APPLICANT:  Rick, Jennifer A.
;  APPLICANT:  Cain, Colyn B.
;  TITLE OF INVENTION: Cancer-linked Genes as Target for Chemotherapy
;  FILE REFERENCE: 689290-273
;  CURRENT APPLICATION NUMBER: US/11/599,845A
;  CURRENT FILING DATE:  2006-11-15
;  PRIOR APPLICATION NUMBER: 10/585,466
;  PRIOR FILING DATE: 2005-01-04
;  PRIOR APPLICATION NUMBER: PCT/US2005/000040
;  PRIOR FILING DATE: 2005-01-04
;  PRIOR APPLICATION NUMBER: 10/583,832
;  PRIOR FILING DATE: 2004-12-16
;  PRIOR APPLICATION NUMBER: PCT/US2004/42406
;  PRIOR FILING DATE: 2004-12-16
;  PRIOR APPLICATION NUMBER: 10/575,337
;  PRIOR FILING DATE: 2004-10-07
;  PRIOR APPLICATION NUMBER: PCT/US2004/33072
;  PRIOR FILING DATE: 2004-10-07
;  PRIOR APPLICATION NUMBER: 10/540,310
;  PRIOR FILING DATE: 2003-12-19
;  PRIOR APPLICATION NUMBER: PCT/US2003/40710
;  PRIOR FILING DATE: 2003-12-19
;  PRIOR APPLICATION NUMBER: 10/518,039
;  PRIOR FILING DATE: 2003-06-10
;  PRIOR APPLICATION NUMBER: PCT/US2003/19741
;  PRIOR FILING DATE: 2003-06-10
;  Remaining Prior Application data removed - See File Wrapper or PALM.
;  NUMBER OF SEQ ID NOS: 769
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 698
;   LENGTH: 843
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-599-845A-698

  Query Match               88.2%;  Score 4364.5;  DB 7;  Length 843;
  Best Local Similarity    99.6%;  Pred. No. 0;
  Matches  824;  Conservative    0;  Mismatches    0;  Indels    3;  Gaps    2;

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Qy        181  YYAEDLRLKLPLQELPNQASNWSAGLLAWLGIPNVLLEVVPDVPPEYYSCRFRVNKLPRF 240
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        182  YYAEDLRLKLPLQELPNQASNWSAGLLAWLGIPNVLLEVVPDVPPEYYSCRFRVNKLPRF 241

Qy        241  LGSDNQDTFFTSTKRHQILFEILAKTPYGHEKKNLLGIHQLLAEGVLSAAFPLHDGPFKT 300
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        242  LGSDNQDTFFTSTKRHQILFEILAKTPYGHEKKNLLGIHQLLAEGVLSAAFPLHDGPFKT 301

Qy        301  PPEGPQAPRLNQRQVLFQHWARWGKWNKYQPLDHVRRYFGEKVALYFAWLGFYTGWLLPA 360
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        302  PPEGPQAPRLNQRQVLFQHWARWGKWNKYQPLDHVRRYFGEKVALYFAWLGFYTGWLLPA 361

Qy        361  AVVGTLVFLVGCFLVFSDIPTQELCGSKDSFEMCPLCLDCPFWLLSSACALAQAGRLFDH 420
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        362  AVVGTLVFLVGCFLVFSDIPTQELCGSKDSFEMCPLCLDCPFWLLSSACALAQAGRLFDH 421

Qy        421  GGTVFFSLFMALWAVLLLEYWKRKSATLAYRWDCSDYEDTEERPRPQFAASAPMTAPNPI 480
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        422  GGTVFFSLFMALWAVLLLEYWKRKSATLAYRWDCSDYEDTEERPRPQFAASAPMTAPNPI 481

Qy        481  TGEDEPYFPERSRARRMLAGSVVIVVMVAVVVMCLVSIILYRAIMAIVVSRSGNTLLAAW 540
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        482  TGEDEPYFPERSRARRMLAGSVVIVVMVAVVVMCLVSIILYRAIMAIVVSRSGNTLLAAW 541

Qy        541  ASRIASLTGSVVNLVFILILSKIYVSLAHVLTRWEMHRTQTKFEDAFTLKVFIFQFVNFY 600
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        542  ASRIASLTGSVVNLVFILILSKIYVSLAHVLTRWEMHRTQTKFEDAFTLKVFIFQFVNFY 601

Qy        601  SSPVYIAFFKGRFVGYPGNYHTLFGVRNEECAAGGCLIELAQELLVIMVGKQVINNMQEV 660
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        602  SSPVYIAFFKGRFVGYPGNYHTLFGVRNEECAAGGCLIELAQELLVIMVGKQVINNMQEV 661

Qy        661  LIPKLKGWWQKFRLRSKKRKAGASAGASQGPWEDDYELVPCEGLFDEYLEMVLQFGFVTI 720
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        662  LIPKLKGWWQKFRLRSKKRKAGASAGASQGPWEDDYELVPCEGLFDEYLEMVLQFGFVTI 721

Qy        721  FVAACPLAPLFALLNNWVEIRLDARKFVCEYRRPVAERAQDIGIWFHILAGLTHLAVISN 780
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        722  FVAACPLAPLFALLNNWVEIRLDARKFVCEYRRPVAERAQDIGIWFHILAGLTHLAVISN 781

Qy        781  AFLLAFSSDFLPRAYYRWTRAHDLRGFLNFTLARAPSSFAAAHNRTC 827
               |||||||||||||||||||||||||||||||||||||||||||||||
Db        782  AFLLAFSSDFLPRAYYRWTRAHDLRGFLNFTLARAPSSFAAAHNRTC 828


RESULT 4
US-10-450-763-45847
; Sequence 45847, Application US/10450763
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; Publication No. US20050196754A1
; GENERAL INFORMATION:
;  APPLICANT: Hyseq, Inc
;  TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
;  FILE REFERENCE: 790CIP3/US
;  CURRENT APPLICATION NUMBER: US/10/450,763
;  CURRENT FILING DATE:  2003-06-11
;  PRIOR APPLICATION NUMBER: PCT/US01/08631
;  PRIOR FILING DATE: 2001-03-30
;  PRIOR APPLICATION NUMBER: 09/540,217
;  PRIOR FILING DATE: 2000-03-31
;  PRIOR APPLICATION NUMBER: 09/649,167
;  PRIOR FILING DATE: 2000-08-23
;  NUMBER OF SEQ ID NOS: 60736
;  SOFTWARE: Custom
; SEQ ID NO 45847
;   LENGTH: 898
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-10-450-763-45847

  Query Match              75.5%;  Score 3736;  DB 5;  Length 898;
  Best Local Similarity    82.3%;  Pred. No. 0;
  Matches  727;  Conservative    4;  Mismatches   16;  Indels  136;  Gaps      6;

Qy       1 MRMAATAWAGLQGPPLPTLCPAVRTGLYCRDQAHAER----------------------- 37
           |||||||||||||||||||||||||||||||||||||
Db       1 MRMAATAWAGLQGPPLPTLCPAVRTGLYCRDQAHAERATDVVLLAPFCQPKTRSHGTCPP 60

Qy      38 -----------------------W---AMTSETS------------------------SG 47
                                  |   |:|  |                          |
Db      61 TERDPRGEGSTEYPGRVDGIQGWGTRALTGWTDRRLLCQACQTLPPRHWFLPGARGWLGG 120

Qy      48 SHCA--------------------RSRMLRRRAQEEDSTVLIDVSPPEAEKRGSYGSTAH 87
           |  ||                    :||||||||||||||||||||||||||||||||||
Db     121 SPCAHGQESLPSQPSPILLRVESVKSRMLRRRAQEEDSTVLIDVSPPEAEKRGSYGSTAH 180

Qy      88 ASEPGGQQAAACRAGSPAKPRIADFVLVWEEDLKLDRQQDSAARDRTDMHRTWRETFLDN 147
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     181 ASEPGGQQAAACRAGSPAKPRIADFVLVWEEDLKLDRQQDSAARDRTDMHRTWRETFLDN 240

Qy     148 LRAAGLCVDQQDVQDGNTTVHYALLSASWAVLCYYAEDLRLKLPLQELPNQASNWSAGLL 207
           |||||||||||||||||||||||||||||||||||||||||||||:|::  |   |  :
Db     241 LRAAGLCVDQQDVQDGNTTVHYALLSASWAVLCYYAEDLRLKLPLQDYPTRPPTGRPACC 300

Qy     208 AWLGIPNVLLEVVPDVPPEYYSCRFRVNKLPRFLGSDNQDTFFTSTKRHQILFEILAKTP 267
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     301 AWLGIPNVLLEVVPDVPPEYYSCRFRVNKLPRFLGSDNQDTFFTSTKRHQILFEILAKTP 360

Qy     268 YGHEKKNLLGIHQLLAEGVLSAAFPLHDGPFKTPPEGPQAPRLNQRQVLFQHWARWGKWN 327
           |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     361 YGHEKKNLLGIHQLLAEGVLSAAFPLHDGPFKTPPEGPQAPRLNQRQVLFQHWARWGKWN 420

Qy     328 KYQPLDHVRRYFGEKVALYFAWLGFYTGWLLPAAVVGTLVFLVGCFLVFSDIPTQELCGS 387
```

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             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    421 KYQPLDHVRRYFGEKVALYFAWLGFYTGWLLPAAVVGTLVFLVGCFLVFSDIPTQELCGS 480

Qy    388 KDSFEMCPLCLDCPFWLLSSACALAQ----AGRLFDHGGTVFFSLFMALWAVLLLEYWKR 443
             ||||||||||||||||||||||||||    ||||||||||||||||||||||||||||||
Db    481 KDSFEMCPLCLDCPFWLLSSACALAQVREEAGRLFDHGGTVFFSLFMALWAVLLLEYWKR 540

Qy    444 KSATLAYRWDCSDYEDTEERPRPQFAASAPMTAPNPITGEDEPYFPERSRARRMLAGSVV 503
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    541 KSATLAYRWDCSDYEDTEERPRPQFAASAPMTAPNPITGEDEPYFPERSRARRMLAGSVV 600

Qy    504 IVVMVAVVVMCLVSIILYRAIMAIVVSRSGNTLLAAWASRIASLTGSVVNLVFILILSKI 563
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    601 IVVMVAVVVMCLVSIILYRAIMAIVVSRSGNTLLAAWASRIASLTGSVVNLVFILILSKI 660

Qy    564 YVSLAHVLTRWEMHRTQTKFEDAFTLKVFIFQFVNFYSSPVYIAFFKGRFVGYPGNYHTL 623
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    661 YVSLAHVLTRWEMHRTQTKFEDAFTLKVFIFQFVNFYSSPVYIAFFKGRFVGYPGNYHTL 720

Qy    624 FGVRNEECAAGGCLIELAQELLVIMVGKQVINNMQEVLIPKLKGWWQKFRLRSKKRKAGA 683
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    721 FGVRNEECAAGGCLIELAQELLVIMVGKQVINNMQEVLIPKLKGWWQKFRLRSKKRKAGA 780

Qy    684 SAGASQGPWEDDYELVPCEGLFDEYLEM-------------------------------- 711
             |||||||||||||||||||||||||||
Db    781 SAGASQGPWEDDYELVPCEGLFDEYLEMGAGFCPNACPELVPELTEPEKARDQPEARSAG 840

Qy    712 -------VLQFGFVTIFVAACPLAPLFALLNNWVEIRLDARKF 747
                    ||||||||||||||||||||||||||||||||||||
Db    841 QDSRPEAVLQFGFVTIFVAACPLAPLFALLNNWVEIRLDARKF 883


RESULT 5
US-10-104-047-2574
; Sequence 2574, Application US/10104047
; Publication No. US20030236392A1
; GENERAL INFORMATION:
;   APPLICANT: HELIX RESEARCH INSTITUTE
;   TITLE OF INVENTION: No. US20030236392A1e1 full length cDNA
;   FILE REFERENCE: H1-A0105
;   CURRENT APPLICATION NUMBER: US/10/104,047
;   CURRENT FILING DATE:  2002-03-25
;   PRIOR APPLICATION NUMBER:
;   PRIOR FILING DATE:
;   NUMBER OF SEQ ID NOS: 4096
;   SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2574
;   LENGTH: 920
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-10-104-047-2574

  Query Match             30.9%;  Score 1531.5;  DB 4;  Length 920;
  Best Local Similarity   37.9%;  Pred. No. 1.9e-138;
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  Matches   360;  Conservative  168;  Mismatches  316;  Indels  105;  Gaps   29;

Qy      44 TSSGSHCARSRMLRRRAQEEDSTVLID----VSPPEAE-----KRGSYGST---AHASEP 91
           :||||     :::       |   :|   :|    ||  :         : :||    || |
Db       4 SSSGITNGKTKVFHPVA--KDVNILFDELEAVSSPCKDDDSLLHPGNLTSTSDDASRLEA 61

Qy      92 GGGQQAAACRAGS-----PAKPRIADFVLVWEEDLKLDRQQDSAARDRTDMHRTWRETFLD 146
           ||:            :        | || |::||: :           :  :|:    || |
Db      62 GGETVPERNKSNGLYFRDGKCRI-DYILVYRK-----------SNPQTEK----REVFER 105

Qy     147 NLRAAGLCVDQQDVQDGNTTVHYALLSASWAVLCYYAEDLRLKLPLQE----LPNQASNW 202
           |:||  ||  ::::      |:  :   |  |  ||  ||| :  :::|  :      || :
Db     106 NIRAEGLQMEKESSLI-NSDIIFVKLHAPWEVLGRYAEQMNVRMPFRRKIYYLPRRYKFM 164

Qy     203 S------AGLLAWLGIPNVLL--EVVPDVPP-EYYSCRFVNKLPRFLGSDNQDTFFTST 253
           |      ||   :   | |     :|      |   |  |  |  ||||:|   |   | |
Db     165 SRIDKQISRLRRWLPKKPMRLDKETLPDLEENDCYTAPFSQQRIHHFI-IHNKETFFNNA 223

Qy     254 KRHQILFEILAKTPYGHEKKNLLGIHQLLAEGVLSAAFPLHDGPFKTPPEGPQAPRLNQR 313
            | :|:  || :  |  | || :|::::|| |   ||||||:| :::         | |
Db     224 TRSRIVHHILQRIKY-EEGKNKIGLNRLLTNGSYEAAFPLHEGSYRSKNSIRTHGAENHR 282

Qy     314 QVLFQHWARWGKWNKYQPLDHVRRYFGEKVALYFAWLGFYTGWLLPAAVVGTLVFLVGCF 373
           :|:: |  |  |  | |||||||| |||||| |||||:| |||  | ||| :|   ||  |
Db     283 HLLYECWASWGVWYKYQPLDLVRRYFGEKIGLYFAWLGWYTGMLFPAAFIGLFVFLYGVT 342

Qy     374 LVFSDIPTQELCGSKDSFEMCPLC-LDCPFWLLSSACALAQAGRLFDHGGTVFFSLFMAL 432
              :    ::|:| :  |     |||:| ||| || :|  |:    |||:||||:|||||:
Db     343 TLDHSQVSKEVCQATDII-MCPVCDKYCPFMRLSDSCVYAKVTHLFDNGATVFFAVFMAV 401

Qy     433 WAVLLLEYWKRKSATLAYRWDCSDYEDTEERPRPQFAAS-APMTAPNPITGEDEPYFPER 491
           || : |||:|||   :|||||  |:|: | | ||  |   |||:| ::| |||::||  |
Db     402 WATVFLEFWKRRRAVIAYDWDLIDWEEEEEEIRPQFEAKYSKKERMNPISGKPEPYQAFT 461

Qy     492 SRARRMLAGSVVIVVMVAVVVMCLVSIILYRAIMAIVVSRSGNTLLA-AWA-----SRIA 545
             :    |:: :      |  |: ||:   :        :|   |  | ||      |::|
Db     462 DKCSRLIVSASGIFFMICVVIAAVFGIVIYRVVTV-------STFAAFKWALIRNNSQVA 514

Qy     546 SLTGSVV--NLVFILILSKIYVSLAHVLTRWEMHRTQTKFEDAFTLKVFIFQVNFYSSP 603
           : | ||:  |    |:|:| :|   :| :||   ||||  | |  ||:::::|::||||:|||||| |
Db     515 T-TGTAVCINFCIIMLLNVLYEKVALLLTNLEQPRTESEWENSFTLKMFLFQFVNLNSST 573

Qy     604 VYIAFFKGRFVGYPGNYHTLFG-VRNECAAGGCLIELAQELLVIMVGKQVINNMQEVLI 662
           |||||  |||| |:|| |   | |  |     |||| | :: ||:|  ||   || |  |
Db     574 FYIAFFLGRFTGHPGAYLRLINRWRLEECHPSGCLIDLCMQMGIIMVLKQTWNNFMELGY 633

Qy     663 PKLKGWWQKFRLRSKKRKAGASAGASQGPWEDDYELVPCE--GLFDEYLEMVLQFGFVTI 720
           |  :: ||  :     |   :::  |        |  || ||  ||||||||||:|||| ||
Db     634 PLIQNWWTR---RKVRQEHGPERKISFPQWEKDYNLQPMNAYGLFDEYLEMILQFGFTTI 690

Qy     721 FVAACPLAPLFALLLNNWVEIRLDARKFVCEYRRPVAERAQDIGIWFHILAGLTHLAVISN 780
           ||||  ||| || || | :|||| |||||||   ::||:|  |:|:|||||: ||  |:|:|
Db     691 FVAAFPLAPLLALLNNIIEIRLDAYKFVTQWRRPLASRAKDIGIWYGILEGIGILSVITN 750
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Qy    781  AFLLAFSSDFLPRAYYRW----------TRAHDLRGFLNFTLA--------------RAP 816
           ||::|  :|||:||  |  :           |:  |::|  :|:
Db    751  AFVIAITSDFIPRLVYAYKYGPCAGQGEAGQKCMVGYVNASLSVFRISDFENRSEPESDG 810


Qy    817  SSFAAAHNRTCRYRAFRDDDGH-----YSQTYWNLLAIRLAFVIVFEHVVFSVGRLLDLL 871
           |  |:     : ||||:||       |: :|::||  ||||:|||||:|| :  |:  |
Db    811  SEFSGTPLKYCRYRDYRDPPHSLVPYGYTLQFWHVLAARLAFIIVFEHLVFCIKHLISYL 870


Qy    872  VPDIPESVEIKVKREYYLAKQALAENEVLFGTNGTKDEQPKGSELSSHW 920
           :||:|: :  :::|| || :: : | |:        |: :  |    :|
Db    871  IPDLPKDLRDRMRREKYLIQEMMYEAELERLQKERKERKKNGKAHHNEW 919
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RESULT 6
US-11-072-512-2574
; Sequence 2574, Application US/11072512
; Publication No. US20060029945A1
; GENERAL INFORMATION:
; APPLICANT: ISOGAI, TAKAO
; APPLICANT: SUGIYAMA, TOMOYASU
; APPLICANT: OTSUKI, TETSUJI
; APPLICANT: WAKAMATSU, AI
; APPLICANT: SATO, HIROYUKI
; APPLICANT: ISHII, SHIZUKO
; APPLICANT: YAMAMOTO, JUN-ICHI
; APPLICANT: ISONO, YUUKO
; APPLICANT: HIO, YURI
; APPLICANT: OTSUKA, KAORU
; APPLICANT: NAGAI, KEIICHI
; APPLICANT: IRIE, RYOTARO
; APPLICANT: TAMECHIKA, ICHIRO
; APPLICANT: SEKI, NAOHIKO
; APPLICANT: YOSHIKAWA, TSUTOMU
; APPLICANT: OTSUKA, MOTOYUKI
; APPLICANT: NAGAHARI, KENJI
; APPLICANT: MASUHO, YASUHIKO
; TITLE OF INVENTION: Novel full length cDNA
; FILE REFERENCE: 084335-0191
; CURRENT APPLICATION NUMBER: US/11/072,512
; CURRENT FILING DATE: 2005-03-07
; PRIOR APPLICATION NUMBER: US 60/350,978
; PRIOR FILING DATE: 2002-01-25
; PRIOR APPLICATION NUMBER: JP 2001-379298
; PRIOR FILING DATE: 2001-11-05
; NUMBER OF SEQ ID NOS: 4096
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2574
;   LENGTH: 920
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-072-512-2574

  Query Match   30.9%; Score 1531.5; DB 6; Length 920;
  Best Local Similarity   37.9%; Pred. No. 1.9e-138;
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```
 Matches    360;  Conservative   168;  Mismatches   316;  Indels  105;  Gaps   29;

Qy      44 TSSGSHCARSRMLRRRAQEEDSTVLID----VSPPEAE-----KRGSYGST---AHASEP  91
           :|||    ::::    |   :|  :|    |   ||   :       : :|:    || |
Db       4 SSSGITNGKTKVFHPVA--KDVNILFDELEAVSSPCKDDDSLLHPGNLTSTSDDASRLEA  61

Qy      92 GGQQAAACRAGS-----PAKPRIADFVLVWEEDLKLDRQQDSAARDRTDMHRTWRETFLD 146
           ||:            :   |  || |::||: :       :  :|:    || |
Db      62 GGETVPERNKSNGLYFRDGKCRI-DYILVYRK-----------SNPQTEK----REVFER 105

Qy     147 NLRAAGLCVDQQDVQDGNTTVHYALLSASWAVLCYYAEDLRLKLPLQE----LPNQASNW 202
           |:|| || ::::      |: : :  | | || ||| : :::|  :     || :
Db     106 NIRAEGLQMEKESSLI-NSDIIFVKLHAPWEVLGRYAEQMNVRMPFRRKIYYLPRRYKFM 164

Qy     203 S------AGLLAWLGIPNVLL--EVVPDVPP-EYYSCRFVNKLPRFLGSDNQDTFFTST  253
           |      || |      : |    :| | : |         | |    | :  |   |
Db     165 SRIDKQISRLRRWLPKKPMRLDKETLPDLEENDCYTAPFSQQRIHHFI-IHNKETFFNNA 223

Qy     254 KRHQILFEILAKTPYGHEKKNLLGIHQLLAEGVLSAAFPLHDGPFKTPPEGPQAPRLNQR 313
           | :|:  || :  | | || :|::::||  |||||:| :::          | |
Db     224 TRSRIVHHILQRIKY-EEGKNKIGLNRLLTNGSYEAAFPLHEGSYRSKNSIRTHGAENHR 282

Qy     314 QVLFQHWARWGKWNKYQPLDHVRRYFGEKVALYFAWLGFYTGWLLPAAVVGTLVFLVGCF 373
           :|:: ||  | || |  |||| ||||::| | ||| ||| :  || :| || | |
Db     283 HLLYECWASWGVWYKYQPLDLVRRYFGEKIGLYFAWLGWYTGMLFPAAFIGLFVFLYGVT 342

Qy     374 LVFSDIPTQELCGSKDSFEMCPLC-LDCPFWLLSSACALAQAGRLFDHGGTVFFSLFMAL 432
           :     ::|:| : |   |||:|  |||  || |: |:   ||:| ||||||:|||||:
Db     343 TLDHSQVSKEVCQATDII-MCPVCDKYCPFMRLSDSCVYAKVTHLFDNGATVFFAVFMAV 401

Qy     433 WAVLLLEYWKRKSATLAYRWDCSDYEDTEERPRPQFAAS-APMTAPNPITGEDEPYFPER 491
           ||:  |||:|:||: :|| || |:|:||  || |    |      :||:::|:  | |
Db     402 WATVFLEFWKRRRAVIAYDWDLIDWEEEEEEIRPQFEAKYSKKERMNPISGKPEPYQAFT 461

Qy     492 SRARRMLAGSVVIVVMVAVVVMCLVSIILYRAIMAIVVSRSGNTLLA-AWA-----SRIA 545
           : |:: :  |   |: ||:  : :::| :      :| | || |::|
Db     462 DKCSRLIVSASGIFFMICVVIAAVFGIVIYRVVTV-------STFAAFKWALIRNNSQVA 514

Qy     546 SLTGSVV--NLVFILILSKIYVSLAHVLTRWEMHRTQTKFEDAFTLKVFIFQFVNFYSSP 603
           : | ||:  |    |::| :|  :|  :|| ||  ||::::|::||||:|::|||||  |
Db     515 T-TGTAVCINFCIIMLLNVLYEKVALLLTNLEQPRTESEWENSFTLKMFLFQFVNLNSST 573

Qy     604 VYIAFFKGRFVGYPGNYHTLFG-VRNEECAAGGCLIELAQELLVIMVGKQVINNMQEVLI 662
           ||||| ||| ::|: | |       |||  |||||| :: :|||  || |    || |:
Db     574 FYIAFFLGRFTGHPGAYLRLINRWRLEECHPSGCLIDLCMQMGIIMVLKQTWNNFMELGY 633

Qy     663 PKLKGWWQKFRLRSKKRKAGASAGASQGPWEDDYELVPCE--GLFDEYLEMVLQFGFVTI 720
           |  :: ||  :  |   :::  |     | || || ||  |||||||||:||||| ||
Db     634 PLIQNWWTR---RKVRQEHGPERKISFPQWEKDYNLQPMNAYGLFDEYLEMILQFGFTTI 690

Qy     721 FVAACPLAPLFALLNNWVEIRLDARKFVCEYRRPVAERAQDIGIWFHILAGLTHLAVISN 780
           |||| |||| | ||||  : |||||   ||:|||  : ||:||||:|| |:| :|:|
Db     691 FVAAFPLAPLLALLNNIIEIRLDAYKFVTQWRRPLASRAKDIGIWYGILEGIGILSVITN 750
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```
Qy      781  AFLLAFSSDFLPRAYYRW----------TRAHDLRGFLNFTLA--------------RAP 816
             ||::|  :|||:|| | :           : |::| :|:
Db      751  AFVIAITSDFIPRLVYAYKYGPCAGQGEAGQKCMVGYVNASLSVFRISDFENRSEPESDG 810


Qy      817  SSFAAAHNRTCRYRAFRDDDGH-----YSQTYWNLLAIRLAFVIVFEHVVFSVGRLLDLL 871
             | |:    : ||||:||        |: :|::|| ||||:|||||:|| : |: |
Db      811  SEFSGTPLKYCYRYDYRDPPHSLVPYGYTLQFWHVLAARLAFIIVFEHLVFCIKHLISYL 870


Qy      872  VPDIPESVEIKVKREYYLAKQALAENEVLFGTNGTKDEQPKGSELSSHW 920
             :||:|: :  :::|| || ::  | |:          |: :  |  : |
Db      871  IPDLPKDLRDRMRREKYLIQEMMYEAELERLQKERKERKKNGKAHHNEW 919



RESULT 7
US-11-582-861-10841
; Sequence 10841, Application US/11582861
; Publication No. US20070099251A1
; GENERAL INFORMATION:
;  APPLICANT: Zhang, Hui
;  APPLICANT:  Aebersold, Rudolf H.
;  TITLE OF INVENTION: TISSUE- SERUM-DERIVED GLYCOPROTEINS
;  TITLE OF INVENTION:  AND METHODS OF THEIR USE
;  FILE REFERENCE: 460092.404
;  CURRENT APPLICATION NUMBER: US/11/582,861
;  CURRENT FILING DATE:  2006-10-17
;  PRIOR APPLICATION NUMBER: US 60/728,044
;  PRIOR FILING DATE: 2005-10-17
;  NUMBER OF SEQ ID NOS: 14918
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 10841
;   LENGTH: 981
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-582-861-10841

  Query Match             30.9%;  Score 1527.5;  DB 6;  Length 981;
  Best Local Similarity   37.8%;  Pred. No. 5e-138;
  Matches  359;  Conservative  168;  Mismatches  317;  Indels  105;  Gaps   29;

Qy       44  TSSGSHCARSRMLRRRAQEEDSTVLID----VSPPEAE------KRGSYGST---AHASEP 91
             :|||    :::    |  :|  :| |    :||| |         |||| |       || |
Db       65  SSSGITNGKTKVFHPVA--KDVNILFDELEAVSSPCKDDDSLLHPGNLTSTSDDASRLEA 122


Qy       92  GGQQAAACRAGS-----PAKPRIADFVLVWEEDLKLDRQQDSAARDRTDMHRTWRETFLD 146
             ||:        :        | || |::||: :             : :|:    || |
Db      123  GGETVPERNKSNGLYFRDGKCRI-DYILVYRK-----------SNPQTEK----REVFER 166


Qy      147  NLRAAGLCVDQQDVQDGNTTVHYALLSASWAVLCYYAEDLRLKLPLQE----LPNQASNW 202
             :|:|| |    :::     |: : :  | | ||| :||| :|||::|    :  ||   |
Db      167  NIRAEGLQMEKESSLI-NSDIIFVKLHAPWEVLGRYAEQMNVRMPFRRKIYYLPRRYKFM 225


Qy      203  S------AGLLAWLGIPNVLL--EVVPDVPP-EYYSCRFRVNKLPRFLGSDNQDTFFTST 253
             |        :    ||   : |  | :||:  : |:    |:     :|  |:::||| :
Db      226  SRIDKQISRFRRWLPKKPMRLDKETLPDLEENDCYTAPFSQQRIHHFI-IHNKETFFNNA 284
```

```
Qy   254 KRHQILFEILAKTPYGHEKKNLLGIHQLLAEGVLSAAFPLHDGPFKTPPEGPQAPRLNQR 313
             | :|:  || :  |  | || :|:::||  |||||:|| :::          | |
Db   285 TRSRIVHHILQRIKY-EEGKNKIGLNRLLTNGSYEAAFPLHEGSYRSKNSIRTHGAENHR 343

Qy   314 QVLFQHWARWGKWNKYQPLDHVRRYFGEKVALYFAWLGFYTGWLLPAAVVGTLVFLVGCF 373
            :|:: || || | |||||| ||||||:|| |||| | ||| :|  |||  |
Db   344 HLLYECWASWGVWYKYQPLDLVRRYFGEKIGLYFAWLGWYTGMLFPAAFIGLFVFLYGVT 403

Qy   374 LVFSDIPTQELCGSKDSFEMCPLC-LDCPFWLLSSACALAQAGRLFDHGGTVFFSLFMAL 432
            :   ::|:| : |   |||:|  |||  || :| |:   |||:| ||||::||:
Db   404 TLDHSQVSKEVCQATDII-MCPVCDKYCPFMRLSDSCVYAKVTHLFDNGATVFFAVFMAV 462

Qy   433 WAVLLLEYWKRKSATLAYRWDCSDYEDTEERPRPQFAAS-APMTAPNPITGEDEPYFPER 491
            || : ||:|||: | :|| ||  |:|: || |||| |  :  |||:|: |||
Db   463 WATVFLEFWKRRAVIAYDWDLIDWEEEEEEIRPQFEAKYSKKERMNPISGKPEPYQAFT 522

Qy   492 SRARRMLAGSVVIVVMVAVVVMCLVSIILYRAIMAIVVSRSGNTLLA-AWA-----SRIA 545
            : |:: :  | |: ||: :  |::|| :    :| | ||    |::|
Db   523 DKCSRLIVSASGIFFMICVVIAAVFGIVIYRVVTV-------STFAAFKWALIRNNSQVA 575

Qy   546 SLTGSVV--NLVFILILSKIYVSLAHVLTRWEMHRTQTKFEDAFTLKVFIFQFVNFYSSP 603
            : ||:  | |  |::|: :|  :| :|| |  ||::::|::||||:|:|||| ||
Db   576 T-TGTAVCINFCIIMLLNVLYEKVALLLTNLEQPRTESEWENSFTLKMFLFQFVNLNSST 634

Qy   604 VYIAFFKGRFVGYPGNYHTLFG-VRNEECAAGGCLIELAQELLVIMVGKQVINNMQEVLI 662
            ||||| ||| || | |   |: |: :|| ||    :: :||| |  ||  |
Db   635 FYIAFFLGRFTGHPGAYLRLINRWRLEECHPSGCLIDLCMQMGIIMVLKQTWNNFMELGY 694

Qy   663 PKLKGWWQKFRLRSKKRKAGASAGASQGPWEDDYELVPCE--GLFDEYLEMVLQFGFVTI 720
            | :: ||  :  |  |::|  |   |  || ||  |||||||||:|||| ||
Db   695 PLIQNWWTR---RKVRQEHGPERKISFPQWEKDYNLQPMNAYGLFDEYLEMILQFGFTTI 751

Qy   721 FVAACPLAPLFALLNNWVEIRLDARKFVCEYRRPVAERAQDIGIWFHILAGLTHLAVISN 780
            |||| ||| |  |: :|||||||| |:|||   ||| |||||||:  || :|| |||
Db   752 FVAAFPLAPLLALLNNIIEIRLDAYKFVTQWRRPLASRAKDIGIWYGILEGIGILSVITN 811

Qy   781 AFLLAFSSDFLPRAYYRW----------TRAHDLRGFLNFTLA--------------RAP 816
            ||::| :|||:|| | :              : |:| :|:
Db   812 AFVIAITSDFIPRLVYAYKYGPCAGQGEAGQKCMVGYVNASLSVFRISDFENRSEPESDG 871

Qy   817 SSFAAAHNRTCRYRAFRDDDGH-----YSQTYWNLLAIRLAFVIVFEHVVFSVGRLLDLL 871
            | |: |||| ||    |||:||       |::|| ||||| |: : :| |
Db   872 SEFSGTPLKYCRYRDYRDPPHSLVPYGYTLQFWHVLAARLAFIIVFEHLVFCIKHLISYL 931

Qy   872 VPDIPESVEIKVKREYYLAKQALAENEVLFGTNGTKDEQPKGSELSSHW 920
            :||:|:: :  ::||  || ::  : |:           |: :     :|
Db   932 IPDLPKDLRDRMRREKYLIQEMMYEAELERLQKERKERKKNGKAHHNEW 980
```

RESULT 8
US-11-582-861-9875
; Sequence 9875, Application US/11582861
; Publication No. US20070099251A1

```
; GENERAL INFORMATION:
;  APPLICANT: Zhang, Hui
;  APPLICANT:  Aebersold, Rudolf H.
;  TITLE OF INVENTION: TISSUE- AND SERUM-DERIVED GLYCOPROTEINS
;  TITLE OF INVENTION:  AND METHODS OF THEIR USE
;  FILE REFERENCE: 460092.404
;  CURRENT APPLICATION NUMBER: US/11/582,861
;  CURRENT FILING DATE:  2006-10-17
;  PRIOR APPLICATION NUMBER: US 60/728,044
;  PRIOR FILING DATE: 2005-10-17
;  NUMBER OF SEQ ID NOS: 14918
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 9875
;   LENGTH: 1046
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-582-861-9875

  Query Match              30.5%;  Score 1511;  DB 6;  Length 1046;
  Best Local Similarity    39.7%;  Pred. No. 2.2e-136;
  Matches  334;  Conservative  156;  Mismatches  264;  Indels   88;  Gaps   25;

Qy     106 KPRIADFVLVWEEDLKLDRQQDSAARDRTDMHRTWRETFLDNLRAAGLCVDQQDVQDGNT 165
           | || ||::||:         :|::   ||| |||| || :::::        :
Db     226 KKRI-DYILVYR--------------KTNIQYDKRNTFEKNLRAEGLMLEKEPA-IASP 268

Qy     166 TVHYALLSASWAVLCYYAEDLRLKLP----------------LQELPNQASNWSAGLLA 208
           :  :   |  | || ||| | ::|             :|   :  || |
Db     269 DIMFIKIHIPWDTLCKYAERLNIRMPFRKKCYYTDGRSKSMGRMQTYFRRIKNWMA---- 324

Qy     209 WLGIPNVL-LEVVPDV-PPEYYSCRFRVNKLPRFLGSDNQDTFFTSTKRHQILFEILAKT 266
            | ||   |:  |:    |:  |:|||:: |::||:: :|
Db     325 --QNPMVLDKSAFPDLEESDCYTGPFSRARIHHFI-INNKDTFFSNATRSRIVYHMLERT 381

Qy     267 PY--GHEKKNLLGIHQLLAEGVLSAAFPLHDGPFKT----PPEGPQAPRLNQRQVLFQHW 320
           |  |   |  :|| :|:  |   |||| |:| :|:       |||    | | :|:: |
Db     382 KYENGISK---VGIRKLINNGSYIAAFPPHEGAYKSSQPIKTHGPQ----NNRHLLYERW 434

Qy     321 ARWGKWNKYQPLDHVRRYFGEKVALYFAWLGFYTGWLLPAAVVGTLVFLVGCFLVFSDIP 380
           |||| | |:|||| :| |||:: |||||:|| |:||||:|| | | | :|
Db     435 ARWGMWYKHQPLDLIRLYFGEKIGLYFAWLGWYTGMLIPAAIVGLCVFFYGLFTMNNSQV 494

Qy     381 TQELCGSKDSFEMCPLC-LDCPFWLLSSACALAQAGRLFDHGGTVFFSLFMALWAVLLLE 439
           :||:|: : : |||||  :|      |: :| ||   |||:||:::||||| : ||
Db     495 SQEICKATEVF-MCPLCDKNCSLQRLNDSCIYAKVTYLFDNGGTVFFAIFMAIWATVFLE 553

Qy     440 YWKRKSATLAYRWDCSDYEDTEERPRPQFAAS-APMTAPNPITGEDEPYFPERSRARRML 498
           :|||: : | | || ::|: || |||| |    | |||||: ||: |   : |:|
Db     554 FWKRRRSILTYTWDLIEWEEEEETLRPQFEAKYYKMEIVNPITGKPEPHQPSSDKVTRLL 613

Qy     499 AGSVVIVVMVAVVVMCLVSIILYR-AIMAIVVSRSGNTLLAAWASRIASLTGSV-VNLVF 556
           |  |:::|:  :  :::|| :|    |  |: :  | - :|:  :| :| :
Db     614 VSVSGIFFMISLVITAVFGVVVYRLVVMEQFASFKWNFIKQYW--QFATSAAAVCINFII 671
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Qy      557  ILILSKIYVSLAHVLTRWEMHRTQTKFEDAFTLKVFIFQFVNFYSSPVYIAFFKGRFVGY  616
             |::|:  |  :|::|| |  |:::::|:|  ||:|:||||| ||  ||||| |||||:
Db      672  IMLLNLAYEKIAYLLTNLEYPRTESEWENSFALKMFLFQFVNLNSSIFYIAFFLGRFVGH  731


Qy      617  PGNYHTLFG-VRNEECAAGGCLIELAQELLVIMVGKQVINNMQEVLIPKLKGWWQKFRLR  675
             || |: ||   | |||  ||||:|  :: |||  ||: ||  |:  | :: || : ::
Db      732  PGKYNKLFDRWRLEECHPSGCLIDLCLQMGVIMFLKQIWNNFMELGYPLIQNWWSRHKI-  790


Qy      676  SKKRKAGASAGASQGPWEDDYELVP--CEGLFDEYLEMVLQFGFVTIFVAACPLAPLFAL  733
             || ||  ||::| : ||||||| ||||||||| ||||||| ||||||||
Db      791  ----KRGIH-DASIPQWENDWNLQPMNLHGLMDEYLEMVLQFGFTTIFVAAFPLAPLLAL  845


Qy      734  LNNNWVEIRLDARKFVCEYRRPVAERAQDIGIWFHILAGLTHLAVISNAFLLAFSSDFLPR  793
             ||| :||||||| ||| ::|||: || ||||| || |: ||||:||::| :||::||
Db      846  LNNIIEIRLDAYKFVTQWRRPLPARATDIGIWLGILEGIGILAVITNAFVIAITSDYIPR  905


Qy      794  AYYRW----------TRAHDLRGFLNFTLARAP-SSFAAAHNRTCRYRAFR-----DDDG  837
                         : |:|::| :|:  : |:||:|: : ::|| ||  : :|
Db      906  FVYEYKYGPCANHVEPSENCLKGYVNNSLSFFDLSELGMGKSGYCRYRDYRGPPWSSKPY  965


Qy      838  HYSQTYWNLLAIRLAFVIVFEHVVFSVGRLLDLLVPDIPESVEIKVKREYYLAKQALAEN  897
             ::  ||::|| |||:|||||||:|||:|||||:|| :   : |:||:|: :  :::||  || :: : |
Db      966  EFTLQYWHILAARLAFIIVFEHLVFGIKSFIAYLIPDVPKGLHDRIRREKYLVQEMMYEA  1025


Qy      898  EV  899
             |:
Db     1026  EL  1027
```

RESULT 9
US-11-443-428A-801153
; Sequence 801153, Application US/11443428A
; Publication No. US20070083334A1
; GENERAL INFORMATION:
; APPLICANT: Mintz, Liat
; APPLICANT: Xie, Hanqing
; APPLICANT: Dahari, Dvir
; APPLICANT: Levanon, Erez
; APPLICANT: Freilich, Shiri
; APPLICANT: Beck, Nili
; APPLICANT: Zhu, Wei-Yong
; APPLICANT: Wasserman, Alon
; APPLICANT: Hermesh, Chen
; APPLICANT: Azar, Idit
; APPLICANT: Bernstein, Jeanne
; TITLE OF INVENTION: METHODS AND SYSTEMS USEFUL FOR ANNOTATING BIOMOLECULAR SEQUENCES
; FILE REFERENCE: 02/23929
; CURRENT APPLICATION NUMBER: US/11/443,428A
; CURRENT FILING DATE: 2006-05-31
; NUMBER OF SEQ ID NOS: 1034312
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 801153
;   LENGTH: 981
;   TYPE: PRT

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;   ORGANISM: Homo sapiens
US-11-443-428A-801153

  Query Match              30.5%;  Score 1508;  DB 6;  Length 981;
  Best Local Similarity    39.7%;  Pred. No. 3.9e-136;
  Matches  334;  Conservative  156;  Mismatches  264;  Indels  88;  Gaps  25;

Qy    106 KPRIADFVLVWEEDLKLDRQQDSAARDRTDMHRTWRETFLDNLRAAGLCVDQQDVQDGNT 165
          | || ::::||:              ::|::    | ||  |||| || :::::     :
Db    161 KRRI-DYILVYR---------------KTNIPYDKRNTFEKNLRAEGLMLEKEPA-IASP 203

Qy    166 TVHYALLSASWAVLCYYAEDLRLKLP----------------LQELPNQASNWSAGLLA 208
          : :  :  |  || ||||  ::::|                 :  || :  |||| ||
Db    204 DIMFIKIHIPWDTLCKYAERLNIRMPFRKKCYYTDGRSKSMGRMQTYFRRIKNWMA---- 259

Qy    209 WLGIPNVL-LEVVPDV-PPEYYSCRFRVNKLPRFLGSDNQDTFFTSTKRHQILFEILAKT 266
          | ||     ||:   : |:    | |:   :: |:  :|:||||:  |  |:|:  :| :|
Db    260 --QNPMVLDKSAFPDLEESDCYTGPFSRARIHHFI-INNKDTFFSNATRSRIVYHMLERT 316

Qy    267 PY--GHEKKNLLGIHQLLAEGVLSAAFPLHDGPFKT----PPEGPQAPRLNQRQVLFQHW 320
          | |    |  :|| :| :      ||| ||  :|: :|:     | ||   ||  || |
Db    317 KYENGISK---VGIRKLINNGSYIAAFFPHEGAYKSSQPIKTHGPQ----NNRHLLYERW 369

Qy    321 ARWGKWNKYQPLDHVRRYFGEKVALYFAWLGFYTGWLLPAAVVGTLVFLVGCFLVFSDIP 380
          |||| | |:|||| :| ||||: |||||:|||||||| :||:|||| | || | || :
Db    370 ARWGMWYKHQPLDLIRLYFGEKIGLYFAWLGWYTGMLIPAAIVGLCVFFYGLFTMNNSQV 429

Qy    381 TQELCGSKDSFEMCPLC-LDCPFWLLSSACALAQAGRLFDHGGTVFFSLFMALWAVLLLE 439
          :||: : : ||||:|:   |:: |:  :|  |:| |   :|||||: ::||||:|:: | |
Db    430 SQEICKATEVF-MCPLCDKNCSLQRLNDSCIYAKVTYLFDNGGTVFFAIFMAIWATVFLE 488

Qy    440 YWKRKSATLAYRWDCSDYEDTEERPRPQFAAS-APMTAPNPITGEDEPYFPERSRARRML 498
          :|||: : | || ::|: ||  |||| :   |      |||||| ||: :  : |:|
Db    489 FWKRRRSILTYTWDLIEWEEEEETLRPQFEAKYYKMEIVNPITGKPEPHQPSSDKVTRLL 548

Qy    499 AGSVVIVVMVAVVVMCLVSIILYR-AIMAIVVSRSGNTLLAAWASRIASLTGSV-VNLVF 556
            |  |:::|:  :   ::||  |:    |   | | |: |  : |:  :| :|  :
Db    549 VSVSGIFFMISLVITAVFGVVVYRLVVMEQFASFKWNFIKQYW--QFATSAAAVCINFII 606

Qy    557 ILILSKIYVSLAHVLTRWEMHRTQTKFEDAFTLKVFIFQFVNFYSSPVYIAFFKGRFVGY 616
          |::|:  | ::|  | | |:::|::| ||:|:  ||:::||  || |::::|| ::  :|
Db    607 IMLLNLAYEKIAYLLTNLEYPRTESEWENSFALKMFLFQFVNLNSSIFYIAFFLGRFVGH 666

Qy    617 PGNYHTLFG-VRNEECAAGGCLIELAQELLVIMVGKQVINNMQEVLIPKLKGWWQKFRLR 675
          || || |    | |||  | |||:| :: ||| ||: |  |: | :: || : : ::
Db    667 PGKYNKLFDRWRLEECHPSGCLIDLCLQMGVIMFLKQIWNNFMELGYPLIQNWWSRHKI- 725

Qy    676 SKKRKAGASAGASQGPWEDDYELVP--CEGLFDEYLEMVLQFGFVTIFVAACPLAPLFAL 733
           || |    ||  ||:| | |  | ||:    |||||||||||||||||  ||||  |
Db    726 ----KRGIH-DASIPQWENDWNLQPMNLHGLMDEYLEMVLQFGFTTIFVAAFPLAPLLAL 780

Qy    734 LNNWVEIRLDARKFVCEYRRPVAERAQDIGIWFHILAGLTHLAVISNAFLLAFSSDFLPR 793
          ||| :|||||||:||| :: ||| |  || ||| ||| |: :||||::::||:|::: ||
Db    781 LNNIIEIRLDAYKFVTQWRRPLPARATDIGIWLGILEGIGILAVITNAFVIATSDYIPR 840
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```
Qy    794 AYYRW----------TRAHDLRGFLNFTLARAP-SSFAAAHNRTCRYRAFR-----DDDG 837
              |  :              :  |:|::|  :|:        |        :   ||||  :|
Db    841 FVYEYKYGPCANHVEPSENCLKGYVNNSLSFFDLSELGMGKSGYCRYRDYRGPPWSSKPY 900

Qy    838 HYSQTYWNLLAIRLAFVIVFEHVVFSVGRLLDLLVPDIPESVEIKVKREYYLAKQALAEN 897
             ::   ||::||  |||||||||:|  :    :   |:||:|:  :   :::|| ||  :: |
Db    901 EFTLQYWHILAARLAFIIVFEHLVFGIKSFIAYLIPDVPKGLHDRIRREKYLVQEMMYEA 960

Qy    898 EV 899
             |:
Db    961 EL 962


RESULT 10
US-11-177-894-7
; Sequence 7, Application US/11177894
; Publication No. US20060040292A1
; GENERAL INFORMATION:
;  APPLICANT: West, et al.
;  TITLE OF INVENTION: Tumor Markers and Uses Thereof
;  FILE REFERENCE: 2002850-0048
;  CURRENT APPLICATION NUMBER: US/11/177,894
;  CURRENT FILING DATE: 2005-07-08
;  NUMBER OF SEQ ID NOS: 29
;  SOFTWARE: PatentIn version 3.2
; SEQ ID NO 7
;  LENGTH: 960
;  TYPE: PRT
;  ORGANISM: Artificial
;  FEATURE:
;  OTHER INFORMATION: Transmembrane protein
US-11-177-894-7

  Query Match             30.1%;  Score 1488;  DB 6;  Length 960;
  Best Local Similarity   37.6%;  Pred. No. 3.4e-134;
  Matches  363;  Conservative  160;  Mismatches  307;  Indels  136;  Gaps   28;

Qy     26 GLYCRDQAHAERWAMT--SETSSGSHCARSRMLRRRAQEEDSTVLIDVSPPEAEKRGSYG 83
             ||| ||         : :    :  ||     :| |  ||  |:             |
Db     52 GLYFRDGRRKVDYILVYHHKRPSG-----NRTLVRRVQHSDTP--------------SGA 92

Qy     84 STAHASEPGGQQAAACRAGSPAKPRIADFVLVWEEDLKLDRQQDSAARDRTDMHRTWRET 143
             :     |   : |:  |||    |          :|  :|          |  ||
Db     93 RSVKQDHPLPGKGASLDAGSGEPP--------------MDYHEDD--------KRFRREE 130

Qy    144 FLDNLRAAGLCVDQQDVQDGNTTVH---YALLSASWAVLCYYAEDLRLKLPLQELPNQAS 200
             :  ||  ||| ::    :| :| :|    :  : | |||  || |:||:| ::: :
Db    131 YEGNLLEAGLELE----RDEDTKIHGVGFVKIHAPWNVLCREAEFLKLKMPTKKMYH--I 184

Qy    201 NWSAGLLAWLGIPNVLLEVVPDVPPEYYSCR----------FRVNKLPRFLGSDNQDTFF 250
             | : |||     | :|| ::    :  | :         |       | |  || :|:||
Db    185 NETRGLLK--KINSVLQKITDPIQPKVAEHRPQTMKRLSYPFSREKQHLFDLSD-KDSFF 241
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Qy    251  TSTKRHQILFEILAKTPYGHEKKNLLGIHQLLAEGVLSAAFPLHDGPFKTPPEGPQAPRL  310
                |  |   |::|||  :|     |   : :||  ||| ||| |::|||||| :       :
Db    242  DSKTRSTIVYEILKRTTCTKAKYS-MGITSLLANGVYAAAYPLHDGDY-----NGENVEF  295

Qy    311  NQRQVLFQHWARWGKWNKYQPLDHVRRYFGEKVALYFAWLGFYTGWLLPAAVVGTLVFLV  370
              | |::|::  |||:|  :  ||||:| ||:|||||: |||||||| ||  |:||::|| :|||
Db    296  NDRKLLYEEWARYGVFYKYQPIDLVRKYFGEKIGLYFAWLGVYTQMLIPASIVGIIVFLY  355

Qy    371  GCFLVFSDIPTQELCGSKDSFEMCPLC-LDCPFWLLSSACALAQAGRLFDHGGTVFFSLF  429
              ||  : :||: |:|   : :  ||||   |:||||  | |||||:||  | :||:: |:||
Db    356  GCATMDENIPSMEMCDQRHNITMCPLCDKTCSYWKMSSACATARASHLFDNPATVFFSVF  415

Qy    430  MALWAVLLLEYWKRKSATLAYRWDCSDYEDTEE----RPRPQFAA-----SAPMTAPNPI  480
              ||||| :|:|||| |   |||| :  :|: ||     || :: |      |   : |
Db    416  MALWAATFMEHWKRKQMRLNYRWDLTGFEEEEEAVKDHPRAEYEARVLEKSLKKESRNKE  475

Qy    481  TGEDEPYFPERSRARRMLAGSVVIVVMVAVVVMCLVSIILYRAIMAIVVSRSGNTLLAAW  540
              | |:     | ||:  |:   |   ::   :: ::||: |  ||:: :|  | :|
Db    476  T--DKVKLTWRDRFPAYLTNLVSIIFMIAVTFAIVLGVIIYRISMAAALAMNSSPSVRSN  533

Qy    541  ASRIASLTGSVVNLVFILILSKIYVSLAHVLTRWEMHRTQTKFEDAFTLKVFIFQFVNFY  600
               :  |    ::|||  ||:|   ::|     :|   ||: |: :|:  ||:    | |: :||| |
Db    534  IRVTVTATAVIINLVVIILLDEVYGCIARWLTKIEVPKTEKSFEERLIFKAFLLKFVNSY  593

Qy    601  SSPVYIAFFKGRFVGYPGNYHTLF-GVRNEECAAGGCLIELAQELLVIMVGKQVI-NNMQ  658
              :    |:|||||||| ||:|   :|    |  |||| |||||:|| :| :||:|||:| ||:
Db    594  TPIFYVAFFKGRFVGRPGDYVYIFRSFRMEECAPGGCLMELCIQLSIIMLGKQLIQNNLF  653

Qy    659  EVLIPKLKGWWQKFRLRSKKRKAGASAGASQGPWEDDYELVPCEGLFDEYLEMVLQFGFV  718
              |: |||:| |    :  :|| :      :|     | ||:| |:||  ||:|||:||||||
Db    654  EIGIPKMKKLIRYLKLKQQSPPDHEECVKRKQRYEVDYNLEPFAGLTPEYMEMIIQFGFV  713

Qy    719  TIFVAACPLAPLFALLNNWVEIRLDARKFVCEYRRPVAERAQDIGIWFHILAGLTHLAVI  778
              |:|||: ||||||||||| |:|||||||:| |  ||||| |:||:||||::||  |:  |||
Db    714  TLFVASFPLAPLFALLNNIIEIRLDAKKFVTELRRPVAVRAKDIGIWYNILRGIGKLAVI  773

Qy    779  SNAFLLAFSSDFLPRA--YYRWTRAHDLRGFLNFTLARAPSSF-------AAAHN-----  824
              |||::|:|:||:     |  |:  :::     ||| |  |||  |          |
Db    774  INAFVISFTSDFIPRLVYLYMYSKNGTMHGFVNHTL----SSFNVSDFQNGTAPNDPLDL  829

Qy    825  ----RTCRYRAFRD---DDGHY--SQTYWNLLAIRLAFVIVFEHVVFSVGRLLDLLVPDI  875
                    : ||| :||    :    |  |: :| | ||||||||||:::   :    :| ::|||
Db    830  GYEVQICRYKDYREPPWSENKYDISKDFWAVLAARLAFVIVFQNLVMFMSDFVDWVIPDI  889

Qy    876  PESVEIKVKREYYLA----------KQALAENEVLFGTNGTKDEQP------------KG  913
              |: :  :: :|  |          ||| |       |||   |              |
Db    890  PKDISQQIHEKVLMVELFMREEQDKQQLL--ETWMEKERQKDEPPCNHHNTKACPDSLG  947

Qy    914  SELSSH  919
              |   ||
Db    948  SPAPSH  953
```

RESULT 11

```
US-11-177-894-11
; Sequence 11, Application US/11177894
; Publication No. US20060040292A1
; GENERAL INFORMATION:
;  APPLICANT: West, et al.
;  TITLE OF INVENTION: Tumor Markers and Uses Thereof
;  FILE REFERENCE: 2002850-0048
;  CURRENT APPLICATION NUMBER: US/11/177,894
;  CURRENT FILING DATE:  2005-07-08
;  NUMBER OF SEQ ID NOS: 29
;  SOFTWARE: PatentIn version 3.2
; SEQ ID NO 11
;   LENGTH: 840
;   TYPE: PRT
;   ORGANISM: Artificial
;   FEATURE:
;   OTHER INFORMATION: Homo sapiens
US-11-177-894-11

  Query Match            29.9%;  Score 1479.5;  DB 6;  Length 840;
  Best Local Similarity  40.0%;  Pred. No. 1.8e-133;
  Matches  340;  Conservative  152;  Mismatches  270;  Indels  89;  Gaps  22;

Qy      135 DMHRTWRETFLDNLRAAGLCVDQQDVQDGNTTVH---YALLSASWAVLCYYAEDLRLKLP 191
                |  |  || :  ||  ||| ::    :| :|| :   :  : | ||| || :|||:|
Db        6 DDKRFRREEYEGNLLEAGLELE----RDEDTKIHGVGFVKIHAPWNVLCREAEFLKLKMP 61

Qy      192 LQEELPNQASNWSAGLLAWLGIPNVLLEVVPDVPPEYYSCR----------FRVNKLPRFL 241
            ::: :   |  |:|||      | :|| ::   : |:    |       |
Db       62 TKKMYH--INETRGLLK--KINSVLQKITDPIQPKVAEHRPQTMKRLSYPFSREKQHLFD 117

Qy      242 GSDNQDTFFTSTKRHQILFEILAKTPYGHEKKNLLGIHQLLAEGVLSAAFPLHDGPFKTP 301
               || :|:||  |   |:||| :|     |  : :|| ||| || :||:|||| :
Db      118 LSD-KDSFFDSKTRSTIVYEILKRTTCTKAKYS-MGITSLLANGVYAAAYPLHDGDY--- 172

Qy      302 PEGPQAPRLNQRQVLFQHWARWGKWNKYQPLDHVRRYFGEKVALYFAWLGFYTGWLLPAA 361
                 :   | |::|:: |||:| :  ||||:| ||:||||: ||||||| ||  |:||:
Db      173 --NGENVEFNDRKLLYEEWARYGVFYKYQPIDLVRKYFGEKIGLYFAWLGVYTQMLIPAS 230

Qy      362 VVGTLVFLVGCFLVFSDIPTQELCGSKDSFEMCPLC-LDCPFWLLSSACALAQAGRLFDH 420
            :|| :||| | :||| |:| :: |  |  | :|||| | : |:|||:|||||:|| | |:
Db      231 IVGIIVFLYGCATMDENIPSMEMCDQRHNITMCPLCDKTCSYWKMSSACATARASHLFDN 290

Qy      421 GGTVFFSLFMALWAVLLLEYWKRKSATLAYRWDCSDYEDTEERPRPQFAA-----SAPMT 475
            ||||| :||||| | : :|:||||   | ||||| | :|: |:|| :: |      |
Db      291 PATVFFSVFMALWAATFMEHWKRKQMRLNYRWDLTGFEEEEDHPRAEYEARVLEKSLKKE 350

Qy      476 APNPITGEDEPYFPERSRARRMLAGSVVIVVMVAVVVMCLVSIILYRAIMAIVVSRSGNT 535
            : | |: |     |  |   :   :| : :  |: :|:||   ||  | :: : :
Db      351 SRNKET--DKVKLTWRDRFPAYLTNLVSIIFMIAVTFAIVLGVIIYRISMAAAALAMNSSP 408

Qy      536 LLAAWASRIASLTGSVVNLVFILILSKIYVSLAHVLTRWEMHRTQTKFEDAFTLKVFIFQ 595
              : :       : |  ::||| |:: ::|  :|  ||: |: :|:  ||:     | |: :
Db      409 SVRSNIRVTVTATAVIINLVVIILLDEVYGCIARWLTKIEVPKTEKSFEERLIFKAFLLK 468
```

```
Qy    596  FVNFYSSPVYIAFFKGRFVGYPGNYHTLF-GVRNEECAAGGCLIELAQELLVIMVGKQVI  654
               |||  |:    |:|||||||||  ||:|   :|    |  ||||  ||||:||   :|  :||:|||:|
Db    469  FVNSYTPIFYVAFFKGRFVGRPGDYVYIFRSFRMEECAPGGCLMELCIQLSIIMLGKQLI  528

Qy    655  -NNMQEVLIPKLKGWWQKFRLRSKKRKAGASAGASQGPWEDDYELVPCEGLFDEYLEMVL  713
               ||: ||  ||:||:|   :  :||:|    : ||:|   |   |:||  ||:|||||  ||:||:
Db    529  QNNLFEIGIPKMKKLIRYLKLKQQSPPDHEECVKRKQRYEVDYNLEPFAGLTPEYMEMII  588

Qy    714  QFGFVTIFVAACPLAPLFALLNNWVEIRLDARKFVCEYRRPVAERAQDIGIWFHILAGLT  773
               |||||||:|||:  |||||||||||  :|||||||:|||  |  |||||  ||:|||||||::||  |:
Db    589  QFGFVTLFVASFPLAPLFALLNNIIEIRLDAKKFVTELRRPVAVRAKDIGIWYNILRGIG  648

Qy    774  HLAVISNAFLLAFSSDFLPRA--YYRWTRAHDLRGFLNFTLARAPSSF-------AAAHN  824
               |||| :||:::|:|||:||   |  :::   :  ||:||  |||       ||     |  |
Db    649  KLAVIIDAFVISFTSDFIPRLVYLVMYSKNGTMHGFVNHTL----SSFNVSDFQNGTAPN  704

Qy    825  ---------RTCRYRAFRD---DDGHY--SQTYWNLLAIRLAFVIVFEHVVFSVGRLLDL  870
                        :  |||: :|:     :   |  |: :|  :||  |||||||||::|   :    :|
Db    705  DPLDLGYEVQICRYKDYREPPWSENKYDISKDFWAVLAARLAFVIVFQNLVMFMSDFVDW  764

Qy    871  LVPDIPESVEIKVKREYYLA----------KQALAENEVLFGTNGTKDEQP---------  911
               ::|||||: :  :: :|  |          ||| |        ||| |
Db    765  VIPDIPKDISQQIHKEKVLMVELFMREEQDKQQLL--ETCMEKERQKDEPPCNHHNTKAC  822

Qy    912  ---KGSELSSH  919
                  ||   ||
Db    823  PDSLGSPAPSH  833
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RESULT 12
US-11-443-428A-774192
; Sequence 774192, Application US/11443428A
; Publication No. US20070083334A1
; GENERAL INFORMATION:
; APPLICANT: Mintz, Liat
; APPLICANT: Xie, Hanqing
; APPLICANT: Dahari, Dvir
; APPLICANT: Levanon, Erez
; APPLICANT: Freilich, Shiri
; APPLICANT: Beck, Nili
; APPLICANT: Zhu, Wei-Yong
; APPLICANT: Wasserman, Alon
; APPLICANT: Hermesh, Chen
; APPLICANT: Azar, Idit
; APPLICANT: Bernstein, Jeanne
; TITLE OF INVENTION: METHODS AND SYSTEMS USEFUL FOR ANNOTATING BIOMOLECULAR SEQUENCES
; FILE REFERENCE: 02/23929
; CURRENT APPLICATION NUMBER: US/11/443,428A
; CURRENT FILING DATE: 2006-05-31
; NUMBER OF SEQ ID NOS: 1034312
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 774192
;   LENGTH: 999

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;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-443-428A-774192

  Query Match              29.6%;  Score 1464;  DB 6;  Length 999;
  Best Local Similarity    37.4%;  Pred. No. 7.7e-132;
  Matches 344;  Conservative 167;  Mismatches 284;  Indels 124;  Gaps   27;

Qy      80 GSYGSTAHASEPGGQQAAACRAGSPAKPRIADFVLVWEEDLKLDRQQDSAARDRTDMHRT 139
            |  |   ||  ||                            |::|     |:: :    :
Db     125 GETGKEPHAGGPG--------------------------DIELG-PLDALEEERKEQ--- 154

Qy     140 WRETFLDNLRAAGLCVDQQDVQDGNTTVHYALLSASWAVLCYYAEDLRLKLPLQELPNQ- 198
            || |   ||  ||| : ::|:: :    :  :  |  ||   || ::|:| ::     :
Db     155 -REEFEHNLMEAGLEL-EKDLENKSQGSIFVRIHAPWQVLAREAEFLKIKVPTKKEMYEI 212

Qy     199 ------ASNWSAGLLAWLGIPNVLLEVVPDVPPEYYSCRFRVNKLP------RFLGSDNQ 246
                  |  :||         |       :|:| | ||:::::  :       :
Db     213 KAGGSIAKKFSAAL------QKLSSHLQPRV-PEHSNNKMKNLSYPFSREKMYLYNIQEK 265

Qy     247 DTFFTSTKRHQILFEILAKTPYGHEKKNLLGIHQLLAEGVLSAAFPLHDGPFKTPPEGPQ 306
            ||||  :   | :|: ||| :|        |  :||: |:|   :  | ::|||  : :| :
Db     266 DTFFDNATRSRIVHEILKRTACS-RANNTMGINSLIANNIYEAAYPLHDGEYDSPEDD-- 322

Qy     307 APRLNQRQVLFQHWARWGKWNKYQPLDHVRRYFGEKVALYFAWLGFYTGWLLPAAVVGTL 366
               :| |::::|  |||:| : |:|:| :|:|||||: |||||||| || :|:|::|:| :
Db     323 ---MNDRKLLYQEWARYGVFYKFQPIDLIRKYFGEKIGLYFAWLGLYTSFLIPSSVIGVI 379

Qy     367 VFLVGCFLVFSDIPTQELCGSKDSFEMCPLC-LDCPFWLLSSACALAQAGRLFDHGGTVF 425
            ||| || : |||||:|:| :::|    :  | |  |    |:  |  ||| ||| :| :|| |||
Db     380 VFLYGCATIEEDIPSREMCDQQNAFTMCPLCDKSCDYWNLSSACGTAQASHLFDNPATVF 439

Qy     426 FSLFMALWAVLLLEYWKRKSATLAYRWDCSDYEDTEER----PRPQFAA----------- 470
            ||:||||||  : || |||   | | |  :  |: ||| |  ||:
Db     440 FSIFMALWATMFLENWKRLQMRLGYFWDLTGIEEEEERAQEHSRPEYETKVREKMLKESN 499

Qy     471 -SAPMTAPNPIT----GEDEPYFPERSRARRMLAGSVVIVVMVAVVVMCLVSIILYRAIM 525
             ||           |    :||      | |:|:     :|| |
Db     500 QSAVQKLETNTTECGDEDDEDKLTWKDRFPGYLMNFASILFMIALTFSIVFGVIVYRITT 559

Qy     526 AIVVSRSGNTLLAAWASRI---ASLTGSVVNLVFILILSKIYVSLAHVLTRWEMHRTQTK 582
            |  :|   |   |  | :      : | ::||  |||| :|| ::|   ||: |: ::|:
Db     560 AAALS-----LNKATRSNVRVTVTATAVIINLVVILILDEIYGAVAKWLTKIEVPKTEQT 614

Qy     583 FEDAFTLKVFIFQFVNFYSSPVYIAFFKGRFVGYPGNYHTLF-GVRNEECAAGGCLIELA 641
            ||:    || |: ||: |  :|||||||||| |    |   |||  ||||  | :  |||||
Db     615 FEERLILKAFLLKFVNAYSPIFYVAFFKGRFVGRPGSYVYVFDGYRMEECAPGGCLMELC 674

Qy     642 QELLVIMVGKQVI-NNMQEVLIPKLKGWWQKFRLRSKKRKAGASAGA-SQGP--WEDDYE 697
             :| :|:||||:| ||: |: :|||    : ||    : :|| : | | |: |    |:||
Db     675 IQLSIIMLGKQLIQNNIFEIGVPKLK---KLFRKLKDETEAGETDSAHSKHPEQWDLDYS 731

Qy     698 LVPCEGLFDEYLEMVLQFGFVTIFVAACPLAPLFALLNNWVEIRLDARKFVCEYRRPVAE 757
            | |   ||  ||::||::|||||:|:|||||:  ||||:||||:|||:|| | ||| |
```

```
Db      732  LEPYTGLTPEYMEMIIQFGFVTLFVASFPLAPVFALLNNVIEVRLDAKKFVTELRRPDAV 791

Qy      758  RAQDIGIWFHILAGLTHLAVISNAFLLAFSSDFLPRAYYRWTRAHD--LRGFLNFTLA-- 813
             : :||||||  ||:|:   :|||||::| :|||:||  |::: :|:  | ||:| ||:
Db      792  RTKDIGIWFDILSGIGKFSVISNAFVIAITSDFIPRLVYQYSYSHNGTLHGFVNHTLSFF 851

Qy      814  ---------RAPSSFAAAHNRTCRYRAFRD-----DDGHYSQTYWNLLAIRLAFVIVFEH 859
                       :  :|      : ||::  :|:     :    :|: || :|: ||||||:|::
Db      852  NVSQLKEGTQPENSQFDQEVQFCRFKDYREPPWAPNPYEFSKQYWFILSARLAFVIIFQN 911

Qy      860  VVFSVGRLLDLLVPDIPESVEIKVKRE------YYLAKQALAENEVLFGTNGTKDEQPKG 913
             :|  :   |:|  ::||||  :  ::|:|        ::|  :     |:|   :       | |
Db      912  LVMFLSVLVDWMIPDIPTDISDQIKKEKSLLVDFFLKE----EHEKLKLMDEPALRSPGG 967

Qy      914  SELSSHWTPFTVPKA-SQL 931
             :  |         :  |   |||
Db      968  GDRSRSRAASSAPSGQSQL 986
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RESULT 13
US-11-097-143-15228
; Sequence 15228, Application US/11097143
; Publication No. US20050208558A1
; GENERAL INFORMATION:
;  APPLICANT: Venter, J. Craig
;  APPLICANT:  et al.
;  TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
;  TITLE OF INVENTION:  ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
;  TITLE OF INVENTION:  DROSOPHILA GENES.
;  FILE REFERENCE: CL000728
;  CURRENT APPLICATION NUMBER: US/11/097,143
;  CURRENT FILING DATE:  2005-04-04
;  PRIOR APPLICATION NUMBER: 60/157,832
;  PRIOR FILING DATE: 1999-10-05
;  PRIOR APPLICATION NUMBER: 60/160,191
;  PRIOR FILING DATE: 1999-10-19
;  PRIOR APPLICATION NUMBER: 60/161,932
;  PRIOR FILING DATE: 1999-10-28
;  PRIOR APPLICATION NUMBER: 60/164,769
;  PRIOR FILING DATE: 1999-11-12
;  PRIOR APPLICATION NUMBER: 60/173,383
;  PRIOR FILING DATE: 1999-12-28
;  PRIOR APPLICATION NUMBER: 60/175,693
;  PRIOR FILING DATE: 2000-01-12
;  PRIOR APPLICATION NUMBER: 60/184,831
;  PRIOR FILING DATE: 2000-02-24
;  PRIOR APPLICATION NUMBER: 60/191,637
;  PRIOR FILING DATE: 2000-03-23
;  NUMBER OF SEQ ID NOS: 43008
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 15228
;   LENGTH: 1219
;   TYPE: PRT
;   ORGANISM: DROSOPHILA
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US-11-097-143-15228

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  Query Match              29.2%;  Score 1445;  DB 6;  Length 1219;
  Best Local Similarity    35.6%;  Pred. No. 7.3e-130;
  Matches  342;  Conservative  165;  Mismatches  332;  Indels  122;  Gaps   27;

Qy      35 AERWAMTSETSSGSHCARSRML------------RRRAQEEDSTVLIDVSPPEAEKRGSY 82
            |:|    :  :   ||   | :|            |:| :   |:|           |||
Db     249 ADRVNQSYEVMESSH---SNVLPDQFGYRQLIPTERKASDTASSV-----------SGSY 294

Qy      83 GSTAHASEP---GGQQAAACRAGSPAKP-------------RIADFVLVW-EEDLKLDRQ 125
            :  ||:   ||:    |     :          |  |  ||  | |||| :  :
Db     295 YGSRKASKSNSLGGESGDERRVSKQDREGLDPESLMFRDGRRKVDMVLAWEEEDLGVMTE 354

Qy     126 QDSAARDRTDMHRTWRETFLDNLRAAGLCVDQQD-VQDGNTTVHYALLSASWAVLCYYAE 184
            :: ||       | :|:::|  || | ||  |    :  |  | :   :  |   ||
Db     355 AEAKRRDN-------RRSFMENLIKEGLEVELEDKSQSFNEKTFFLKIHLPWRLETRLAE 407

Qy     185 DLRLKLP------------------LQELPNQASNWSAGLLAWLGIPNVLLEVVPDVPP 225
            :  ||||                  |: :     |    |   : :    |||
Db     408 VMNLKLPVKRFITISVKPSWDEENVVLRNMQYWKDVWQR-LTKKIQLDQTLLE----GET 462

Qy     226 EYYSCRFRVNKLPRFLGSDNQDTFFTSTKRHQILFEILAKTPYGHEKKNLLGIHQLLAEG 285
             : :      |  :|:  | | |||:|| :::|  ::| :||:       ::   || :|
Db     463 TFKAATANGNPEEQFIVKD-RATAFTSAQRSLMVMQVLIRTPFDESDRS--GIRRLMNDG 519

Qy     286 VLSAAFPLHDGPFKTPPEGPQAPRLN-QRQVLFQHWARWGKWNKYQPLDHVRRYFGEKVA 344
              ||||:| :    : | :  ::  |:||:| ||  :| |||  |:|||:|:|
Db     520 TYLGCFPLHEGRY----DRPHSSGISLDRRVLYQTWAHPSQWYKKQPLCLVRKYFGDKIA 575

Qy     345 LYFAWLGFYTGWLLPAAVVGTLVFLVGCFLVFSD--IPTQELCG--SKDSFEMCPLC-LD 399
            ||| |||| :  ||||||| |: ||:||| |: : : |     |  |:: |:||| : |
Db     576 LYFCWLGFYTEMLVYPAVVGTLCFIYGLATLESEDNTPSKEICNEYGTGNITLCPLCDKA 635

Qy     400 CPFWLLSSACALAQAGRLFDHGGTVFFSLFMALWAVLLLEYWKRKSATLAYRWDCSDYED 459
            | :  ||  :|  ::   |||: ||||::||: ||   || |||| : | :|    : |
Db     636 CSYQRLSESCLFSRLTYLFDNPSTVFFAIFMSFWATTFLELWKRKQSVLVWEWDLHNV-D 694

Qy     460 TEERPRPQFAASAPMTAPNPITGEDEPYFPERSRARRMLAGSVVIVVMVAVVVMCLVSII 519
             :| ||:|      ||:| |    :|: |  :|:    ::|::||  ::  | |
Db     695 MDEENRPEFETNATTFRMNPVTREKEPYMSTWNRSIRFVITGSAVLFMISVVLSAVLGTI 754

Qy     520 LYRAIMAIVVSRSGNTLLAAWASRIASLTGSVVNLVFILILSKIYVSLAHVLTRWEMHRT 579
            |||   : |:   | :|    :: :|::|::| |:|:|:||::| |  |     |
Db     755 LYRITLVSVIYGGGGFFVKEHAKLFTSVTAALINLVVIMILTRIYHRMAIKLTNLENPRT 814

Qy     580 QTKFEDAFTLKVFIFQFVNFYSSPVYIAFFKGRFVGYPGNYHT----LFGVRNEECAAGG 635
             |::||::|:  |:| |:|:||||  |||||||| :|||||||    |  | ::|:  |  |
Db     815 HTEYEDSYTFKIFFFEFMNFYSSLIYIAFFKGRFFDYPGDDQARKSEFFRLKNDICDPAG 874

Qy     636 CLIELAQELLVIMVGKQVINNMQEVLIPKLKGWWQKFRLRSKKRKAGASAGASQGPWEDD 695
            || ||   :|  :||||||    | |  | |       | :   : ::|         || |
Db     875 CLSELCIQLAIIMVGKQCWNNFMEYLFPKFWNWWR----QRKHKQATKDESHLHMAWEQD 930
```

```
Qy       696 YELV-PCE-GLFDEYLEMVLQFGFVTIFVAACPLAPLFALLNNWVEIRLDARKFVCEYRR 753
             |  :   |    |||||||||:||:||||:||||  ||||||||||| ||  |  |: ||
Db       931 YHMQDPGRLALFDEYLEMILQYGFVTLFVAAFPLAPLFALLNNVAEIRLDAYKMVTQARR 990


Qy       754 PVAERAQDIGIWFHILAGLTHLAVISNAFLLAFSSDFLPRAYYR--WTRAHDLRGFLNFT 811
             |:|||  :||| |:  ||   :|: ||:|||||:|::::|||:|  |:   ::   ||:: :
Db       991 PLAERVEDIGAWYGILRIITYTAVVSNAFVIAYTSDFIPRMVYKFVYSETHTLAGYIEHS 1050


Qy       812 LA---------RAPSSFAAAHNRTCRYRAFRDDDGHY-----SQTYWNLLAIRLAFVIVF 857
             |:          :|: |||:|| :|: |   |  ||::|   | ||||||||
Db      1051 LSIFNTSDYKEEWGASVSEKDPDTCQYRGYRNGPKDYEPYGLSPHYWHVFAARLAFVVVF 1110


Qy       858 EHVVFSVGRLLDLLVPDIPESVEIKVKREYYLAKQALAENEVLFGTNGTKDEQPKGSELS 917
             |||||  :  ::   ::||:|  |: :::||   |||:|   :   :|  |    ::
Db      1111 EHVVFVITGIMQFIIPDVPSEVKTQMQREQLLAKEAKYQ-------HGIKRAQGDSQDIM 1163


Qy       918 S 918
             |
Db      1164 S 1164
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RESULT 14
US-11-443-428A-739452
; Sequence 739452, Application US/11443428A
; Publication No. US20070083334A1
; GENERAL INFORMATION:
; APPLICANT: Mintz, Liat
; APPLICANT: Xie, Hanqing
; APPLICANT: Dahari, Dvir
; APPLICANT: Levanon, Erez
; APPLICANT: Freilich, Shiri
; APPLICANT: Beck, Nili
; APPLICANT: Zhu, Wei-Yong
; APPLICANT: Wasserman, Alon
; APPLICANT: Hermesh, Chen
; APPLICANT: Azar, Idit
; APPLICANT: Bernstein, Jeanne
; TITLE OF INVENTION: METHODS AND SYSTEMS USEFUL FOR ANNOTATING BIOMOLECULAR SEQUENCES
; FILE REFERENCE: 02/23929
; CURRENT APPLICATION NUMBER: US/11/443,428A
; CURRENT FILING DATE: 2006-05-31
; NUMBER OF SEQ ID NOS: 1034312
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 739452
;   LENGTH: 800
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-443-428A-739452

  Query Match   28.6%; Score 1417; DB 6; Length 800;
  Best Local Similarity   40.3%; Pred. No. 2e-127;
  Matches 326; Conservative 142; Mismatches 252; Indels   88; Gaps   22;

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Qy       179 LCYYAEDLRLKLPLQELPNQASNWSAGLLAWLGIPNVLLEVVPDVPPEYYSCR------- 231
```

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                ||   |   |:||:|  :::  :     |  :  |||     |  :||  ::    :  |:    |
Db        7  LC--ARVLKLKMPTKKMYH--INETRGLLK--KINSVLQKITDPIQPKVAEHRPQTMKRL  60

Qy      232  ---FRVNKLPRFLGSDNQDTFFTSTKRHQILFEILAKTPYGHEKKNLLGIHQLLAEGVLS  288
                |   |  ||  |:|:||  |  |||||  :|       |   |  |:|||  |||| |
Db       61  SYPFSREKQHLFDLSD-KDSFFDSKTRSTIVYEILKRTTCTKAKYS-MGITSLLANGVYA  118

Qy      289  AAFPLHDGPFKTPPEGPQAPRLNQRQVLFQHWARWGKWNKYQPLDHVRRYFGEKVALYFA  348
             ||:||||| :        :      |  |::|::  |||:|  :  ||||:|  ||:|||||   ||||
Db      119  AAYPLHDGDY-----NGENVEFNDRKLLYEEWARYGVFYKYQPIDLVRKYFGEKIGLYFA  173

Qy      349  WLGFYTGWLLPAAVVGTLVFLVGCFLVFSDIPTQELCGSKDSFEMCPLC-LDCPFWLLSS  407
             |||  ||  ||||| :  ||  |   :  :||: |:  :  |||||    | :| :||
Db      174  WLGVYTQMLIPASIVGIIVFLYGCATMDENIPSMEMCDQRHNITMCPLCDKTCSYWKMSS  233

Qy      408  ACALAQAGRLFDHGGTVFFSLFMALWAVLLLEYWKRKSATLAYRWDCSDYEDTEE----R  463
             ||| |:|   |||:   |||||:|||||||       :|:||||    |  ||||  :  :|:  ||
Db      234  ACATARASHLFDNPATVFFSVFMALWAATFMEHWKRKQMRLNYRWDLTGFEEEEEAVKDH  293

Qy      464  PRPQFAA-----SAPMTAPNPITGEDEPYFPERSRARRMLAGSVVIVVMVAVVVMCLVSI  518
             ||  ::   |       :  |   |  | :| |  :  |||  ||||
Db      294  PRAEYEARVLEKSLKKESRNKET--DKVKLTWRDRFPAYLTNLVSIIFMIAVTFAIVLGV  351

Qy      519  ILYRAIMAIVVSRSGNTLLAAWASRIASLTGSVVNLVFILILSKIYVSLAHVLTRWEMHR  578
             |:||  ||  :: : :   : :      :  | :|||  |:|| ::|   :|    |||  :: :
Db      352  IIYRISMAAALAMNSSPSVRSNIRVTVTATAVIINLVVIILLDEVYGCIARWLTKIEVPK  411

Qy      579  TQTKFEDAFTLKVFIFQFVNFYSSPVYIAFFKGRFVGYPGNYHTLF-GVRNEECAAGGCL  637
             |:   ||:   |  |: :||  |  |||  |||||||  :|    ||  |  :|   |||
Db      412  TEKSFEERLIFKAFLLKFVNSYTPIFYVAFFKGRFVGRPGDYVVIFRSFRMEECAPGGCL  471

Qy      638  IELAQELLVIMVGKQVI-NNMQEVLIPKLKGWWQKFRLRSKKRKAGASAGASQGPWEDDY  696
             :||   :|  :||:|||:|  |||  |: |  |||:|    :  :|: :               :  :|  ||
Db      472  MELCIQLSIIMLGKQLIQNNLFEIGIPKMKKLIRYLKLKQQSPPDHEECVKRKQRYEVDY  531

Qy      697  ELVPCEGLFDEYLEMVLQFGFVTIFVAACPLAPLFALLNNWVEIRLDARKFVCEYRRPVA  756
             |  |   ||   ||:||::|||||:||| ||||||||||||| :||||||||:||||||:|||  |  |||||
Db      532  NLEPFAGLTPEYMEMIIQFGFVTLFVASFPLAPLFALLNNIIEIRLDAKKFVTELRRPVA  591

Qy      757  ERAQDIGIWFHILAGLTHLAVISNAFLLAFSSDFLPRA--YYRWTRAHDLRGFLNFTLAR  814
             ||:||||||   :||  |||:    ||::|||   ||:|::     |   :::  ||   |||
Db      592  VRAKDIGIWYNILRGIGKLAVIINAFVISFTSDFIPRLVYLYMYSKNGTMHGFVNHTL--  649

Qy      815  APSSF-------AAAHN---------RTCRYRAFRD---DDGHY--SQTYWNLLAIRLAF  853
                |||       |  |         :  |||: :|:   :   |  |:  :|  ||||
Db      650  --SSFNVSDFQNGTAPNDPLDLGYEVQICRYKDYREPPWSENKYDISKDFWAVLAARLAF  707

Qy      854  VIVFEHVVFSVGRLLDLLVPDIPESVEIKVKREYYLA----------KQALAENEVLFGT  903
             |||||:|||   ||| :  :::|||:||  :  :| :             ||  |    |||
Db      708  VIVFQNLVFMFSDFVDWVIPDIPKDISQQIHKEKVLMVELFMREEQDKQQLL--ETWMEK  765

Qy      904  NGTKDEQP------------KGSELSSH  919
                |||            ||   |||
Db      766  ERQKDEPPCNHHNTKACPDSLGSPAPSH  793
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RESULT 15
US-11-443-428A-739454
; Sequence 739454, Application US/11443428A
; Publication No. US20070083334A1
; GENERAL INFORMATION:
;  APPLICANT: Mintz, Liat
;  APPLICANT:  Xie, Hanqing
;  APPLICANT:  Dahari, Dvir
;  APPLICANT:  Levanon, Erez
;  APPLICANT:  Freilich, Shiri
;  APPLICANT:  Beck, Nili
;  APPLICANT:  Zhu, Wei-Yong
;  APPLICANT:  Wasserman, Alon
;  APPLICANT:  Hermesh, Chen
;  APPLICANT:  Azar, Idit
;  APPLICANT:  Bernstein, Jeanne
;  TITLE OF INVENTION: METHODS AND SYSTEMS USEFUL FOR ANNOTATING BIOMOLECULAR SEQUENCES
;  FILE REFERENCE: 02/23929
;  CURRENT APPLICATION NUMBER: US/11/443,428A
;  CURRENT FILING DATE:  2006-05-31
;  NUMBER OF SEQ ID NOS: 1034312
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 739454
;   LENGTH: 800
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-11-443-428A-739454

  Query Match            28.6%;  Score 1417;  DB 6;  Length 800;
  Best Local Similarity  40.3%;  Pred. No. 2e-127;
  Matches  326;  Conservative  142;  Mismatches  252;  Indels   88;  Gaps   22;

Qy      179 LCYYAEDLRLKLPLQELPNQASNWSAGLLAWLGIPNVLLEVVPDVPPEYYSCR------- 231
                ||   |   |:||:| ::: :    | : |||    | :|| ::    : |:    |
Db        7 LC--ARVLKLKMPTKKMYH--INETRGLLK--KINSVLQKITDPIQPKVAEHRPQTMKRL 60

Qy      232 ---FRVNKLPRFLGSDNQDTFFTSTKRHQILFEILAKTPYGHEKKNLLGIHQLLAEGVLS 288
               |   |   || :|:|| |   | |:::||| :|     | : :|| ||| || :
Db       61 SYPFSREKQHLFDLSD-KDSFFDSKTRSTIVYEILKRTTCTKAKYS-MGITSLLANGVYA 118

Qy      289 AAFPLHDGPFKTPPEGPQAPRLNQRQVLFQHWARWGKWNKYQPLDHVRRYFGEKVALYFA 348
               ||:||||| :         :     |  |::|:: |||:| : |||:| ||:|||||: ||||
Db      119 AAYPLHDGDY-----NGENVEFNDRKLLYEEWARYGVFYKYQPIDLVRKYFGEKIGLYFA 173

Qy      349 WLGFYTGWLLPAAVVGTLVFLVGCFLVFSDIPTQELCGSKDSFEMCPLC-LDCPFWLLSS 407
               ||| ||  |:||::||  :||| ||   :  :||: |:|   : :  ||||||   | :| :||
Db      174 WLGVYTQMLIPASIVGIIVFLYGCATMDENIPSMEMCDQRHNITMCPLCDKTCSYWKMSS 233

Qy      408 ACALAQAGRLFDHGGTVFFSLFMALWAVLLLEYWKRKSATLAYRWDCSDYEDTEE----R 463
               ||| |:| |||: ||||:||||||| :|:|||| | |||| : :|: ||
Db      234 ACATARASHLFDNPATVFFSVFMALWAATFMEHWKRKQMRLNYRWDLTGFEEEEEAVKDH 293
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Qy      464 PRPQFAA-----SAPMTAPNPITGEDEPYFPERSRARRMLAGSVVIVVMVAVVVMCLVSI 518
                |  ::  |       :  |  |:      |  |       |  |  |:|  |    ::  :
Db      294 PRAEYEARVLEKSLKKESRNKET--DKVKLTWRDRFPAYLTNLVSIIFMIAVTFAIVLGV 351

Qy      519 ILYRAIMAIVVSRSGNTLLAAWASRIASLTGSVVNLVFILILSKIYVSLAHVLTRWEMHR 578
                |:||   ||   ::  :   :  :      :  |   ::|||  ||:|  ::|    :|   ||:  |: :
Db      352 IIYRISMAAALAMNSSPSVRSNIRVTVTATAVIINLVVIILLDEVYGCIARWLTKIEVPK 411

Qy      579 TQTKFEDAFTLKVFIFQFVNFYSSPVYIAFFKGRFVGYPGNYHTLF-GVRNEECAAGGCL 637
                |:   ||  :|  :|||  :||| ||    |:|||||||||  ||:|    :|      |||| |||
Db      412 TEKSFEERLIFKAFLLKFVNSYTPIFYVAFFKGRFVGRPGDYVYIFRSFRMEECAPGGCL 471

Qy      638 IELAQELLVIMVGKQVI-NNMQEVLIPKLKGWWQKFRLRSKKRKAGASAGASQGPWEDDY 696
                :||   :|  :||:|||:|  ||:  |:  |||:|    :   :|:  :             :  :|  ||
Db      472 MELCIQLSIIMLGKQLIQNNLFEIGIPKMKKLIRYLKLKQQSPPDHEECVKRKQRYEVDY 531

Qy      697 ELVPCEGLFDEYLEMVLQFGFVTIFVAACPLAPLFALLNNWVEIRLDARKFVCEYRRPVA 756
                 |  |   ||:|::||||||:|||:  |||||||||  :|||||:||| :||||||| |
Db      532 NLEPFAGLTPEYMEMIIQFGFVTLFVASFPLAPLFALLNNIIEIRLDAKKFVTELRRPVA 591

Qy      757 ERAQDIGIWFHILAGLTHLAVISNAFLLAFSSDFLPRA--YYRWTRAHDLRGFLNFTLAR 814
                ||:||||::||  |:    ||||  |||::::|:|||:||      |  :::     :  ||:|  ||
Db      592 VRAKDIGIWYNILRGIGKLAVIINAFVISFTSDFIPRLVYLYMYSKNGTMHGFVNHTL-- 649

Qy      815 APSSF-------AAAHN---------RTCRYRAFRD---DDGHY--SQTYWNLLAIRLAF 853
                 |||       |  |       :  |||:  :|:     :  |  |: :|  :||  ||||
Db      650 --SSFNVSDFQNGTAPNDPLDLGYEVQICRYKDYREPPWSENKYDISKDFWAVLAARLAF 707

Qy      854 VIVFEHVVFSVGRLLDLLVPDIPESVEIKVKREYYLA----------KQALAENEVLFGT 903
                ||||::::|   :|  ::||||:  :  ::  |  |            ||  |    |
Db      708 VIVFQNLVMFMSDFVDWVIPDIPKDISQQIHKEKVLMVELFMREEQDKQQLL--ETWMEK 765

Qy      904 NGTKDEQP------------KGSELSSH 919
                 |||  |                   ||   ||
Db      766 ERQKDEPPCNHHNTKACPDSLGSPAPSH 793
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Search completed: October 1, 2008, 13:00:05
Job time : 830 secs